OIPE

RAW SEQUENCE LISTING   DATE: 11/27/2001
PATENT APPLICATION: US/09/989,481   TIME: 14:18:41

Input Set : A:\Chau, Raymond 12592-4 Sequence Listing in PatentIn 3.0.txt
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 3  <110> APPLICANT: Chau, Raymond M.W.
 5  <120> TITLE OF INVENTION: Isolation and Use of Motoneurontropic Factors
 7  <130> FILE REFERENCE: 12592-4
 9  <140> CURRENT APPLICATION NUMBER: US/09/989,481
10  <141> CURRENT FILING DATE: 2001-11-20
12  <150> PRIOR APPLICATION NUMBER: US 09/633,447
13  <151> PRIOR FILING DATE: 2000-08-07
15  <150> PRIOR APPLICATION NUMBER: US 08/9928862
16  <151> PRIOR FILING DATE: 1997-09-12
18  <150> PRIOR APPLICATION NUMBER: US 08/751225
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21  <150> PRIOR APPLICATION NUMBER: US 60/026792
22  <151> PRIOR FILING DATE: 1996-09-27
24  <160> NUMBER OF SEQ ID NOS: 5
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38  gacacctttg gtgacattaa ttaccaagaa tttgctaaaa gactctgggg tgacatctac  180
40  ttcaacccta agacgcgaaa gttcaccaaa aaggccccaa ctagcagctc ccagagaagt  240
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44  gacaccagcc tcccacggac cctagacgag cttggcatcc acctgacgaa ggaggagctg  360
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52  gaccctgatg gccccctgat gtgccacact actaagatgt tcagcacaca tgatggagtc  600
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56  gttctggggg agaactacac cctggaggat gaggaagact ccccaatttg ccccgtgggc  720
58  cgcctttgga tctctgtggc cagctaccac atcgaggtga accgtgttcc tgctggcaac  780
60  tgggttctga ttgaaggtgt tgatcaacca attgtgaaga cagcaaccat aaccgaaccc  840
62  cgaggcaatg aggaggctca gattttccga cccttgaagt tcaataccac atctgttatc  900
64  aagattgctg tggagccagt caacccctca gagctgccca agatgcttga tggcctgcgc  960
66  aaggtcaaca agagctatcc atccctcacc accaaggtgg aggagtctgg cgagcatgtg 1020
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72  acatcctccc tcaagtgctt tgctgaaacg cctaataaga gaacaagat caccatgatt 1200
74  gctgagcctc ttgagaaggg cctggcagag gacatagaga atgaggtggt ccagattacg 1260
76  tggaacagga agaagctggg agagttcttc cagaccaagt acgattggga tctgctggct 1320
78  gcccgttcca tctgggcttt tggccctgat gcgactggcc ccaacattct ggtggatgat 1380
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82  ggt                                                             1443
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86  <211> LENGTH: 927
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88  <213> ORGANISM: Homo Sapiens
90  <400> SEQUENCE: 2
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93  gctcgatgtc ttgcagaagg acatgatggt cctacacagt aaggaatgga ttacctacaa   120
95  tattaatagc agcctcccat acacactttt gacacccttc cctaaaggat taatatgctc   180
97  caaccttcct gtccccacag ttcagtggct ctccctaccc tcaccatgat cggatgaaaa   240
99  aaaataaggt ttcacagctt aagagtgaaa ttctggaatc caactacaag ctcataactg   300
101 tagcatggaa cctggtagta gcataataaa taaatttta gtaagaggct taagaaattt    360
103 tagcaaaaaa agcactccct ttcttcctcc ctacatatct catatgtttt tcaacacaaa   420
105 aaattctgtg attttagaga aacttcttac agtactttta agttcaaaac cagatgctca   480
107 ttacagttct tttaaacacc aaactagtca tctcaaaaat atggctaact ctctggacta   540
109 aattccatag gaaaaattat taatttcaaa atgcctaatt tttgatcaat gctgaagagc   600
111 caagcaatca tgtcctgctt ctcactcagg gcagagttct caggtcagaa gctccggagt   660
113 ctgtcagaga ttaaaatatc atctcaacaa ttcacaagct acttctaagt gttaccctaa   720
115 attagtcact aatcgtttct cccccaactc tatttcacaa attaaagttt acagaattga   780
117 caaaaaccaa accaatgaaa caacccaggc tatttgcagg gggggggaaa gagatacccc   840
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134 Gly Asn Val Cys Phe Ser Ser Ser Gln Tyr Ser Ile Cys Phe Thr Leu
135             20                  25                  30
137 Gly Ser Phe Ala Lys Ile Tyr Ala Asp Thr Phe Gly Asp Ile Asn Tyr
138         35                  40                  45
140 Gln Glu Phe Ala Lys Arg Leu Trp Gly Asp Ile Tyr Phe Asn Pro Lys
141     50                  55                  60
143 Thr Arg Lys Phe Thr Lys Lys Ala Pro Thr Ser Ser Ser Gln Arg Ser
144 65                  70                  75                  80
146 Phe Val Glu Phe Ile Leu Glu Pro Leu Tyr Lys Ile Leu Ala Gln Val
147             85                  90                  95
149 Val Gly Asp Val Asp Thr Ser Leu Pro Arg Thr Leu Asp Glu Leu Gly
150             100                 105                 110
152 Ile His Leu Thr Lys Glu Glu Leu Lys Leu Asn Ile Arg Pro Leu Leu
153         115                 120                 125
155 Arg Leu Val Cys Lys Lys Phe Phe Gly Glu Phe Thr Gly Phe Val Asp
156     130                 135                 140
158 Met Cys Val Gln His Ile Pro Ser Pro Lys Val Gly Ala Lys Pro Lys
159 145                 150                 155                 160
161 Ile Glu His Thr Tyr Thr Gly Gly Val Asp Ser Asp Leu Gly Glu Ala
162             165                 170                 175
164 Met Ser Asp Cys Asp Pro Asp Gly Pro Leu Met Cys His Thr Thr Lys
165             180                 185                 190
167 Met Phe Ser Thr His Asp Gly Val Gln Phe His Pro Phe Gly Arg Val
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168           195                200                205
170 Leu Ser Gly Thr Ile His Ala Gly Gln Pro Val Lys Val Leu Gly Glu
171     210                215                220
173 Asn Tyr Thr Leu Glu Asp Glu Glu Asp Ser Gln Ile Cys Thr Val Gly
174 225                230                235                240
176 Arg Leu Trp Ile Ser Val Ala Arg Tyr His Ile Glu Val Asn Arg Val
177             245                250                255
179 Pro Ala Gly Asn Trp Val Leu Ile Glu Gly Val Asp Gln Pro Ile Val
180             260                265                270
182 Lys Thr Ala Thr Ile Thr Glu Pro Arg Gly Asn Glu Glu Ala Gln Ile
183     275                280                285
185 Phe Arg Pro Leu Lys Phe Asn Thr Thr Ser Val Ile Lys Ile Ala Val
186     290                295                300
188 Glu Pro Val Asn Pro Ser Glu Leu Pro Lys Met Leu Asp Gly Leu Arg
189 305                310                315                320
191 Lys Val Asn Lys Ser Tyr Pro Ser Leu Thr Thr Lys Val Glu Glu Ser
192             325                330                335
194 Gly Glu His Val Ile Leu Gly Thr Gly Glu Leu Tyr Leu Asp Cys Val
195             340                345                350
197 Met His Asp Leu Arg Lys Met Tyr Ser Glu Ile Asp Ile Lys Val Ala
198     355                360                365
200 Asp Pro Val Val Thr Phe Cys Glu Thr Val Val Glu Thr Ser Ser Leu
201     370                375                380
203 Lys Cys Phe Ala Glu Thr Pro Asn Lys Lys Asn Lys Ile Thr Met Ile
204 385                390                395                400
206 Ala Glu Pro Leu Glu Lys Gly Leu Ala Glu Asp Ile Glu Asn Glu Val
207             405                410                415
209 Val Gln Ile Thr Trp Asn Arg Lys Lys Leu Gly Glu Phe Phe Gln Thr
210     420                425                430
212 Lys Tyr Asp Trp Asp Leu Leu Ala Ala Arg Ser Ile Trp Ala Phe Gly
213     435                440                445
215 Pro Asp Ala Thr Gly Pro Asn Ile Leu Val Asp Asp Thr Leu Pro Ser
216     450                455                460
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219 465                470                475                480
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225 <211> LENGTH: 33
226 <212> TYPE: PRT
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229 <400> SEQUENCE: 4
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235             20                25                30
237 Gln
240 <210> SEQ ID NO: 5
241 <211> LENGTH: 99
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245 <400> SEQUENCE: 5
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248 gctcgatgtc ttgcagaagg acatgatggt cctacacag   99

VERIFICATION SUMMARY   DATE: 11/27/2001
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Input Set : **A:\Chau, Raymond 12592-4 Sequence Listing in PatentIn 3.0.txt**
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L:9 M:270 C: Current Application Number differs, Replaced Current Application Number